Protein sequences of th   examples (mutated to yield soluble forms)

>sFcγRI   SEQ ID NO: 1

MAVISLQPPWVSVFQEETVTLHCEvLHLPGSSSTQWFLNGTATQTSTPSYRITSASVNDS
GEYRCQRGLSGRSDPIQLEIHRGWLLLQVSSRVFTEGEPLALRCHAWKDKLVYNVLYYRN
GKAFKFFHWNSNLTILKTNISHNGTYHCSGMGKHRYTSAGISVTVKELFPAPVLNASVTS
PLLEGNLVTLSCETKLLLQRPGLQLYFSFYMGSKTLRGRNTSSEYQILTARREDSGLYWC
EAATEDGNVLKRSPELELQVLGLQLPTPV

                                 SEQ ID NO: 2

>sFcγRIIa

MAAPPKAVLKLEPPWINVLQEDSVTLTCQGARSPESDSIQWFHNGNLIPTHTQPSYRFKA
NNNDSGEYTCQTGQTSLSDPVHLTVLSEWLVLQTPHLEFQEGETIMLRCHSWKDKPLVKV
TFFQNGKSQKFSRLDPTFSIPQANHSHSGDYHCTGNIGYTLFSSKPVTITVQVP

                                 SEQ ID NO: 3

· ₃FcγRIIb

..uGTPAAPPKAVLKLEPQWINVLQEDSVTLTCRGTHSPESDSIQWFHNGNLIPTHTQPSYR
FKANNNDSGEYTCQTGQTSLSDPVHLTVLSEWLVLQTPHLEFQEGETIVLRCHSWKDKPL
VKVTFFQNGKSKKFSRSDPNFSIPQANHSHSGDYHCTGNIGYTLYSSKPVTITVQAPSSS
PMGII

                                 SEQ ID NO: 4

>sFcγRIII

MRTEDLPKAVVFLEPQWYSVLEKDSVTLKCQGAYSPEDNSTQWFHNESLISSQASSYFID
AATVNDSGEYRCQTNLSTLSDPVQLEVHIGWLLLQAPRWVFKEEDPIHLRCHSWKNTALH
KVTYLQNGKDRKYFHHNSDFHIPKATLKDSGSYFCRGLVGSKNVSSETVNITITQG

                                 SEQ ID NO: 5

>sFcεRIa

MAVPQKPKVSLNPPWNRIFKGENVTLTCNGNNFFEVSSTKWFHNGSLSEETNSSLNIVN
AKFEDSGEYKCQHQQVNESEPVYLEVFSDWLLLQASAEVVMEGQPLFLRCHGWRNWDVYK
VIYYKDGEALKYWYENHNISITNATVEDSGTYYCTGKVWQLDYESEPLNITVIKAPREKY
ʷˡLQF

                                 SEQ ID NO: 6

>sFcεRII

MDTTQSLKQLEERAARNVSQVSKNLESHHGDQMTQKSQSTQISQELEELRAEQQRLKSQD
LELSWNLNGLQADLSSFKSQELNERNEASDLLERLREEVTKLRMELQVSSGFVCNTCPEK
WINFQRKCYYFGKGTKQWVHARYACDDMEGQLVSIHSPEEQDFLTKHASHTGSWIGLRNL
DLKGEFIWVDGSHVDYSNWAPGEPTSRSQGEDCVMMRGSGRWNDAFCDRKLGAWVCDRLA
TCTPPASEGSAESMGPDSRPDPDGRLPTPSAPLHS

DNA sequences of the examples (mutated to yield soluble forms)

>sFγRI   SEQ ID NO: 7
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  1 CATATGGCAG TGATCTCTTT GCAGCCTCCA TGGGTCAGCG TGTTCCAAGA GGAAACCGTA   60
 61 ACCTTGCACT GTGAGGTGCT CCATCTGCCT GGGAGCAGCT CTACACAGTG GTTTCTCAAT  120
121 GGCACAGCCA CTCAGACCTC GACCCCCAGC TACAGAATCA CCTCTGCCAG TGTCAATGAC  180
181 AGTGGTGAAT ACAGGTGCCA GAGAGGTCTC TCAGGGCGAA GTGACCCCAT ACAGCTGGAA  240
241 ATCCACAGAG GCTGGCTACT ACTGCAGGTC TCCAGCAGAG TCTTCACGGA AGGAGAACCT  300
301 CTGGCCTTGA GGTGTCATGC GTGGAAGGAT AAGCTGGTGT ACAATGTGCT TTACTATCGA  360
361 AATGGCAAAG CCTTTAAGTT TTTCCACTGG AATTCTAACC TCACCATTCT GAAAACCAAC  420
421 ATAAGTCACA ATGGCACCTA CCATTGCTCA GGCATGGGAA AGCATCGCTA CACATCAGCA  480
481 GGAATATCTG TCACTGTGAA AGAGCTATTT CCAGCTCCAG TGCTGAATGC ATCTGTGACA  540
541 TCCCCACTCC TGGAGGGGAA TCTGGTCACC CTGAGCTGTG AAACAAAGTT GCTCTTGCAG  600
601 AGGCCTGGTT TGCAGCTTTA CTTCTCCTTC TACATGGGCA GCAAGACCCT GCGAGGCAGG  660
661 AACACATCCT CTGAATACCA AATAGTAACT GCTAGAAGAG AAGACTCTGG GTTATACTGG  720
721 TGCGAGGCTG CCACAGAGGA TGGAAATGTC CTTAAGCGCA GCCCTGAGTT GGAGCTTCAA  780
781 GTGCTTGGCC TCCAGTTACC AACTCCTGTC TAGTCTCGAG                         820
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>sFcγRIIa   SEQ ID NO: 8
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  1 CATATGGCAG CTCCCCCAAA GGCTGTGCTG AAACTTGAGC CCCCGTGGAT CAACGTGCTC   60
 61 CAGGAGGACT CTGTGACTCT GACATGCCAG GGGGCTGCCA GCCCTGAGAG CGACTCCATT  120
121 CAGTGGTTCC ACAATGGGAA TCTCATTCCC ACCCACACGC AGCCCAGCTA CAGGTTCAAG  180
181 GCCAACAACA ATGACAGCGG GGAGTACACG TGCCAGACTG GCCAGACCAG CCTCAGCGAC  240
241 CCTGTGCATC TGACTGTGCT TTCCGAATGG CTGGTGCTCC AGACCCCTCA CCTGGAGTTC  300
301 CAGGAGGGAG AAACCATCAT GCTGAGGTGC CACAGCTGGA AGGACAAGCC TCTGGTCAAG  360
361 GTCACATTCT TCCAGAATGG AAAATCCCAG AAATTCTCCC GTTTGGATCC CACCTTCTCC  420
421 ATCCCACAAG CAAACCACAG TCACAGTGGT GATTACCACT GCACAGGAAA CATAGGCTAC  480
481 ACGCTGTTCT CATCCAAGCC TGTGACCATC ACTGTCCAAG TGCCCTGAAG CTT          533
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>sFcγRIIb   SEQ ID NO: 9
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  1 CCATGGGGAC ACCTGCAGCT CCCCCAAAGG CTGTGCTGAA ACTCGAGCCC CAGTGGATCA   60
 61 ACGTGCTCCA GGAGGACTCT GTGACTCTGA CATGCCGGGG GACTCACAGC CCTGAGAGCG  120
121 ACTCCATTCA GTGGTTCCAC AATGGGAATC TCATTCCCAC CCACACGCAG CCCAGCTACA  180
181 GGTTCAAGGC CAACAACAAT GACAGCGGGG AGTACACGTG CCAGACTGGC CAGACCAGCC  240
241 TCAGCGACCC TGTGCATCTG ACTGTGCTTT CTGAGTGGCT GGTGCTCCAG ACCCCTCACC  300
301 TGGAGTTCCA GGAGGGAGAA ACCATCGTGC TGAGGTGCCA CAGCTGGAAG GACAAGCCTC  360
361 TGGTCAAGGT CACATTCTTC CAGAATGGAA AATCCAGAAG ATTTTCCCGT TCGGATCCCA  420
421 ACTTCTCCAT CCCACAAGCA AACCACAGTC ACAGTGGTGA TTACCACTGC ACAGGAAACA  480
481 TAGGCTACAC GCTGTACTCA TCCAAGCCTG TGACCATCAC TGTCCAAGCT CCCAGCTCTT  540
541 CACCGATGGG GATCATTTAG GCTGTCGAC                                     569
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>sFcγRIII   SEQ ID NO: 10
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  1 CATATGCGGA CTGAAGATCT CCCAAAGGCT GTGGTGTTCC TGGAGCCTCA ATGGTACAGC   60
 61 GTGCTTGAGA AGGACAGTGT GACTCTGAAG TGCCAGGGAG CCTACTCCCC TGAGGACAAT  120
121 TCCACACAGT GGTTTCACAA TGAGAGCCTC ATCTCAAGCC AGGCCTCGAG CTACTTCATT  180
181 GACGCTGCCA CAGTCAACGA CAGTGGAGAG TACAGGTGCC AGACAAACCT CTCCACCCTC  240
241 AGTGACCCGG TGCAGCTAGA AGTCCATATC GGCTGGCTGT TGCTCCAGGC CCCTCGGTGG  300
301 GTGTTCAAGG AGGAAGACCC TATTCACCTG AGGTGTCACA GCTGGAAGAA CACTGCTCTG  360
361 CATAAGGTCA CATATTTACA GAATGGCAAA GACAGGAAGT ATTTTCATCA TAATTCTGAC  420
421 TTCCACATTC CAAAAGCCAC ACTCAAAGAT AGCGGCTCCT ACTTCTGCAG GGGGCTTGTT  480
481 GGGAGTAAAA ATGTGTCTTC AGAGACTGTG AACATCACCA TCACTCAAGG TTAAGCTT     538
```

>sFcεRIa   SEQ ID NO: 11
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  1 CATATGGCAG TCCCTCAGAA ACCTAAGGTC TCCTTGAACC CTCCATGGAA TAGAATATTT   60
 61 AAAGGAGAGA ATGTGACTCT TACATGTAAT GGGAACAATT TCTTTGAAGT CAGTTCCACC   120
121 AAATGGTTCC ACAATGGCAG CCTTTCAGAA GAGACAAATT CAAGTTTGAA TATTGTGAAT  180
181 GCCAAATTTG AAGACAGTGG AGAATACAAA TGTCAGCACC AACAAGTTAA TGAGAGTGAA  240
241 CCTGTGTACC TGGAAGTCTT CAGTGACTGG CTGCTCCTTC AGGCCTCTGC TGAGGTGGTG  300
301 ATGGAGGGCC AGCCCCTCTT CCTCAGGTGC CATGGTTGGA GGAACTGGGA TGTGTACAAG  360
361 GTGATCTATT ATAAGGATGG TGAAGCTCTC AAGTACTGGT ATGAGAACCA CAACATCTCC  420
421 ATTACAAATG CCACAGTTGA AGACAGTGGA ACCTACTACT GTACGGGCAA AGTGTGGCAG  480
481 CTGGACTATG AGTCTGAGCC CCTCAACATT ACTGTAATAA AAGCTCCGCG TGAGAAGTAC  540
541 TGGCTACAAT TTTAGGATCC                                               560
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09/856933

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                                    SEQ ID NO: 12
>sFcɛRII
    1 CATATGGAGT TGCAGGTGTC CAGCGGCTTT GTGTGCAACA CGTGCCCTGA AAAGTGGATC    60
   61 AATTTCCAAC GGAAGTGCTA CTACTTCGGC AAGGGCACCA AGCAGTGGGT CCACGCCCGG   120
  121 TATGCCTGTG ACGACATGGA AGGGCAGCTG GTCAGCATCC ACAGCCCGGA GGAGCAGGAC   180
  181 TTCCTGACCA AGCATGCCAG CCACACCGGC TCCTGGATTG GCCTTCGGAA CTTGGACCTG   240
  241 AAGGGGGAGT TTATCTGGGT GGATGGGAGC CACGTGGACT ACAGCAACTG GGCTCCAGGG   300
  301 GAGCCCACCA GCCGGAGCCA GGGCGAGGAC TGCGTGATGA TGCGGGGCTC CGGTCGCTGG   360
  361 AACGACGCCT TCTGCGACCG TAAGCTGGGC GCCTGGGTGT GCGACCGGCT GGCCACATGC   420
  421 ACGCCGCCAG CCAGCGAAGG TTCCGCGGAG TCCATGGGAC CTGATTCAAG ACCAGACCCT   480
  481 GACGGCCGCC TGCCCACCCC CTCTGCCCCT CTCCACTCTT GAGCATGGAT CC            532
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SEQ ID NO:13

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human FcγRIIb2
   1    ggctgtgact gctgtgctct gggcgccact cgctccaggg agtgatggga atcctgtcat
   61   ttttacctgt ccttgccact gagagtgact gggctgactg caagtccccc cagccttggg
  121   gtcatatgct tctgtggaca gctgtgctat tcctggctcc tgttgctggg acacctgcag
  181   ctcccccaaa ggctgtgctg aaactcgagc cccagtggat caacgtgctc caggaggact
  241   ctgtgactct gacatgccgg gggactcaca gccctgagag cgactccatt cagtggttcc
  301   acaatgggaa tctcattccc acccacacgc agcccagcta caggttcaag gccaacaaca
  361   atgacagcgg ggagtacacg tgccagactg gccagaccag cctcagcgac cctgtgtcatc
  421   tgacagtgct ttctgagtgg ctggtgctcc agacccctca cctggagttc caggagggag
  481   aaaccatcgt gctgaggtgc cacagctgga ggacaagcct ctggtcaag gtcacattct
  541   tccagaatgg aaaatccaag aaattttccc gttcggatcc caacttctcc atcccacaag
  601   caaaccacag tcacagtggt gattaccatt gcacaggaaa cataggctac acgctgtact
  661   catccaagcc tgtgaccatc actgtccaag ctcccagctc ttcaccgatg gggatcattg
  721   tggctgtggt cactgggatt gctgtagctg ccattgttgc tgctgtagtg gccttgatct
  781   actgcaggaa aaagcggatt tcagccaatc ccactaatcc tgatgaggct gacaaagttg
  841   gggctgagaa cacaatcacc tattcacttc tcatgcaccc ggatgctctg gaagagcctg
  901   atgaccagaa ccgtatttag tctccattgt cttgcattgg gatttgagaa gaaatcagag
  961   agggaagatc tggtatttcc tggcctaaat tccccttggg gaggacaggg agatgctgca
 1021   gttccaaaag agaaggtttc ttccagagtc atctacctga atcctgaagc tccctgtcct
 1081   gaaagccaca gacaatatgg tcccaaatgc ccgactgcac cttctgtgct tcagctcttc
 1141   ttgacatcaa ggctcttccg ttccacatcc acacagccaa tccaattaat caaaccactg
 1201   ttattaacag ataatagcaa cttgggaaat gcttatgtta caggttacgt gagaacaatc
 1261   atgtaaatct atatgatttc agaaatgtta aaatagacta acctctacca gcacattaaa
 1321   agtgattgtt tctgggtgat aaaattattg atgattttta ttttctttat ttttctataa
 1381   agatcatata ttacttttat aataaaacat tataaaaac
```

SEQ ID NO:14

human FcεRIa

```
1      agatctcagc acagtaagca ccaggagtcc atgaagaaga tggctcctgc catggaatcc
61     cctactctac tgtgtgtagc cttactgttc ttcgctccag atggcgtgtt agcagtccct
121    cagaaaccta aggtctcctt gaaccctcca tggaatagaa tatttaaagg agagaatgtg
181    actcttacat gtaatgggaa caatttcttt gaagtcagtt ccaccaaatg gttccacaat
241    ggcagccttt cagaagagac aaattcaagt ttgaatattg tgaatgccaa atttgaagac
301    agtggagaat acaaatgtca gcaccaacaa gttaatgaga gtgaacctgt gtacctggaa
361    gtcttcagtg actggctgct ccttcaggcc tctgctgagg tggtgatgga gggccagccc
421    ctcttcctca ggtgccatgg ttggaggaac tgggatgtgt acaaggtgat ctattataag
481    gatggtgaag ctctcaagta ctggtatgag aaccacaaca tctccattac aaatgccaca
541    gttgaagaca gtggaaccta ctactgtacg ggcaaagtgt ggcagctgga ctatgagtct
601    gagcccctca acattactgt aataaaagct ccgcgtgaga agtactggct acaatttttt
661    atcccattgt tggtggtgat tctgtttgct gtggacacag gattatttat ctcaactcag
721    cagcaggtca catttctctt gaagattaag agaaccagga aaggcttcag acttctgaac
781    ccacatccta agccaaaccc caaaaacaac tgatataatt aactcaagaa atatttgcaa
841    cattagtttt tttccagcat cagcaattgc tactcaattg tcaaacacag cttgcaatat
901    acatagaaac gtctgtgtctc aaggatttat agaaatgctt cattaaactg agtgaaactg
961    attaagtggc atgtaatagt aagtgctcaa ttaacattgg ttgaataaat gagagaatga
1021   atagattcat ttattagcat ttgtaaaaga gatgttcaat ttagatct
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SEQ ID NO:15

human mRNA for high affinity Fc receptor (FcγRI)

```
1     gacagatttc actgctccca ccagcttgga gacaacatgt ggttcttgac aactctgctc
61    ctttgggttc cagttgatgg gcaagtggac accacaaagg cagtgatctc tttgcagcct
121   ccatgggtca gcgtgttcca agaggaaacc gtaaccttgc actgtgaggt gctccatctg
181   cctgggagca gctctacaca gtggtttctc aatggcacag ccactcagac ctcgaccccc
241   agctacagaa tcacctctgc cagtgtcaat gacagtggtg aatacaggtg ccagagaggt
301   ctctcagggc gaagtgaccc catacagctg gaaatccaca gaggctggct actactgcag
361   gtctccagca gagtcttcac ggaaggagaa cctctggcct tgaggtgtca tgcgtggaag
421   gataagctga tgtacaatgt gctttactat cgaaatggca aagccttaa gtttttccac
481   tggaattcta acctcaccat tctgaaaacc aacataagtc acaatggcac ctaccattgc
541   tcaggcatgg gaaagcatcg ctacacatca gcaggaatat ctgtcactgt gaaagagcta
601   tttccagctc cagtgctgaa tgcatctgtg acatccccac tcctggaggg gaatctggtc
661   accctgagct gtgaaacaaa gttgctcttg cagaggcctg gtttgcagct ttacttctcc
721   ttctacatgg gcagcaagac cctgcgaggc aggaacacat cctctgaata ccaaatacta
781   actgctagaa gagaagactc tgggttatac tggtgcgagg ctgccacaga ggatggaaat
841   gtccttaagc gcagccctga gttggagctt caagtgcttg gcctccagtt accaactcct
901   gtctggtttc atgtcctttt ctatctggca gtgggaataa tgttttagt gaacactgtt
961   ctctgggtga caatacgtaa agaactgaaa agaaagaaa agtgggattt agaaatctct
1021  ttggattctg gtcatgagaa gaagtaact tccagccttc aagaagacag acatttagaa
1081  gaagagctga aatgtcagga acaaaaagaa gaacagctgc aggaaggggt gcaccggaag
1141  gagccccagg gggccacgta gcagcggctc agtgggtggc catcgatctg gaccgtcccc
1201  tgcccacttg ctccccgtga gcactgcgta caaacatcca aaagttcaac aacaccagaa
1261  ctgtgtgtct catggtatgt aactcttaaa gcaaataaat gaactgactt caaaaaaaaa
1321  a
```

SEQ ID NO:16

human FcγRIIa

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1     cccaaatgtc tcagaatgta tgtcccagaa acctgtggct gcttcaacca ttgacagttt
61    tgctgctgct ggcttctgca gacagtcaag ctgcagctcc cccaaaggct gtgctgaaac
121   ttgagcccc gtggatcaac gtgctccagg aggactctgt gactcgtgaca tgccagggg
181   ctcgcagccc tgagagcgac tccattcagt ggttccacaa tgggaatctc attcccaccc
241   acacgcagcc cagctacagg ttcaaggcca acaacaatga cagcgggag tacacgtgcc
301   agactggcca gaccagcctc agcgacctg tgcatctgac tgtgctttcc gaatggctgg
361   tgctccagac ccctcacctg gagttccagg aggggagaaac catcatgctg aggtgccaca
421   gctggaagga caagcctctg gtcaaggtca cattcttcca gaatggaaaa tcccagaaat
481   tctcccgttt ggatccccac ttctcccatcc cacaagcaaa ccacagtcac agtggtgatt
541   accactgcac aggaaacata ggctacacgc tgttctcatc caagcctgtg accatcactg
601   tccaagtgcc cagcatgggc agctcttcac caatggggat cattgtggct gtggtcattg
661   cgactgctgt agcagccatt gttgctgctg tagtggcctt gatctactgc aggaaaaagc
721   ggatttcagc caattccact gatcctgtga ggctgtccca atttgagcca cctggacgtc
781   aaatgattgc catcagaaag agacaacttg aagaaaccaa caatgactat gaaacagctg
841   acggcggcta catgactctg aaccccaggg cacctactga cgatgataaa aacatctacc
901   tgactcttcc tcccaacgac catgtcaaca gtaataacta aagagtaacg ttatgccagg
961   tggtcatact ctcagcttgc tgatggatga caaaaagagg ggaattgtta aaggaaaatt
1021  taaatggaga ctggaaaaat cctgagcaa caaaccacc tggccttag aaatagcttt
1081  aactttgctt aaactacaaa cacaagcaaa acttcacggg gtcatactac atacaagcat
1141  aagcaaaact taacttggat catttctggt aaatgcttat gttagaaata agacaacccc
1201  agccaatcac aagcagccta ctaacatata attaggtgac tagggacttt ctaagaagat
1261  acctacccc aaaaaacaat tatgtaattg aaaaccaacc gattgccttt attttgcttc
1321  cacattttcc caataaatac ttgcctgtga cattttgcca ctggaacact aaacttcatg
1381  aattgcgcct cagattttc ctttaacatc tttttttttt ttgacagagt ctcaatctgt
1441  tacccaggct ggagtgcagt ggtgctatct tggctcactg caaacccgcc tcccaggttt
1501  aagcgattct tatgcctcag cctcccagta gctgggatta gaggcatgtg ccatcatacc
1561  cagctaattt ttgtattttt tattttttat ttttagtaga gacagggttt cgcaatgttg
1621  gccaggccga tctcgaactt ctggcctcta gcgatctgcc cgcctcggcc tcccaaagtg
1681  ctgggatgac cgcatcagcc ccaatgtcca gcctctttaa catcttcttt cctatgccct
1741  ctctgtggat ccctactgct ggttvctgcc ttctccatgc tgagaacaaa atcacctatt
1801  cactgcttat gcagtcggaa gctccagaag aacaaagagc ccaattacca gaaccacatt
1861  aagtctccat tgtttgcct tgggatttga gaagagaatt agagaggtga ggatctggta
1921  tttcctggac taaattccct tggggaagac gaagggatgc tgcagttcca aaagagaagg
1981  actcttccag agtcatctac ctgagtccca aagctccctg tcctgaaagc cacagacaat
2041  atggtcccaa atgactgact gcaccttctg tgcctcagcc gttcttgaca tcaagaatct
2101  tctgttccac atccacacag ccaatacaat tagtcaaacc actgttatta acagatgtag
2161  caacatgaga aacgcttatg ttacaggtta catgagagca atcatgtaag tctatatgac
2221  ttcagaaatg ttaaaataga ctaacctcta acaacaaatt aaaagtgatt gtttcaaggt
2281  gatgcaatta ttgatgacct attttatttt tctataatga tcatatatta cctttgtaat
2341  aaaacattat aaccaaaac
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SEQ ID NO:17

human FcγRIII

```
1     tctttggtga cttgtccact ccagtgtggc atcatgtggc agctgctcct cccaactgct
61    ctgctacttc tagtttcagc tggcatgcgg actgaagatc tcccaaaggc tgtggtgttc
121   ctggagcctc aatggtacag cgtgcttgag aaggacagtg tgactctgaa gtgccaggga
181   gcctactccc ctgaggacaa ttccacacag tggtttcaca atgagagcct catctcaagc
241   caggcctcga gctacttcat tgacgctgcc acagtcaacg acagtggaga gtacaggtgc
301   cagacaaacc tctccaccct cagtgacccg gtgcagctag aagtccatat cggctggctg
361   ttgctccagg cccctcggtg ggtgttcaag gaggaagacc ctattcacct gaggtgtcac
421   agctggaaga acactgctct gcataaggtc acatatttac agaatggcaa agacaggaag
481   tattttcatc ataattctga cttccacatt ccaaaagcca cactcaaaga tagcggctcc
541   tacttctgca gggggcttgt tgggagtaaa aatgtgtctt cagagactgt gaacatcacc
601   atcactcaag gtttggcagt gtcaaccatc tcatcattct ctccacctgg gtaccaagtc
661   tctttctgct tggtgatggt actccttttt gcagtggaca caggactata tttctctgtg
721   aagacaaaca tttgaagctc aacaagagac tggaaggacc ataaacttaa atggagaaag
781   gaccctcaag acaaatgacc cccatcccat gggagtaata agagcagtgg cagcagcatc
841   tctgaacatt tctctggatt tgcaaccccca tcatcctcag gcctctc
```

SEQ ID NO: 18

human FcεRII

```
        1 ctcctgctta aacctctgtc tctgacggtc cctgccaatc gctctggtcg
accccaacac
       61 actaggagga cagacacagg cttcaaactc cactaagtga ccagagctgt gattgtgccc
      121 gctgagtgga ctgcgttgtc agggagtgag tgctccatca tcgggagaat ccaagcagga
      181 ccgccatgga ggaaggtcaa tattcagaga tcgaggagct tcccaggagg cggtgttgca
      241 ggcgtgggac tcagatcgtg ctgctggggc tggtgaccgc cgctctgtgg gctgggctgc
      301 tgactctgct tctcctgtgg cactgggaca ccacacagag tctaaaacag ctggaagaga
      361 gggctgcccg gaacgtctct caagtttcca agaacttgga aagccaccac ggtgaccaga
      421 tggcgcagaa atcccagtcc acgcagattt cacaggaact ggaggaactt cgagctgaac
      481 agcagagatt gaaatctcag gacttggagc tgtcctggaa cctgaacggg cttcaagcag
      541 atctgagcag cttcaagtcc caggaattga acgagaggaa cgaagcttca gatttgctgg
      601 aaagactccg ggaggaggtg acaaagctaa ggatggagtt gcaggtgtcc agcggctttg
      661 tgtgcaacac gtgccctgaa aagtggatca atttccaacg gaagtgctac tacttcggca
      721 agggcaccaa gcagtgggtc cacgcccggt atgcctgtga cgacatggaa gggcagctgg
      781 tcagcatcca cagccccgag gagcaggact tcctgaccaa gcatgccagc cacaccggct
      841 cctggattgg ccttcggaac ttggacctga agggagagtt tatctgggtg gatgggagcc
      901 atgtggacta cagcaactgg gctccagggg agcccaccag ccggagccag ggcgaggact
      961 gcgtgatgat gcggggctcc ggtcgctgga acgacgcctt ctgcgaccgt aagctgggcg
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     1201 ggcctaaaag cctctttgtg gctgaaaggt ccctgtgaca ttttctgcca cccaaacgga
     1261 ggcagctgac acattcccg ctcctctatg gccctgcctc tcccaggagt acacccaac
     1321 agcaccctct ccagatggga gtgcccccaa cagcaccctc tccagatgag agtacacccc
     1381 aacagcaccc tctccagatg cagccccatc tcctcagcac cccaggacct gagtatcccc
     1441 agctcaggtg gtgagtcctc ctgtccagcc tgcatcaata aaatgggca gtgatggcct
     1501 ccc
```